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Date: 07/27/99

## LARGE-SCALE CsCl preparation of bacterial genomic DNA

1. Grow 100 ml culture of bacterial strain to saturation.
2. Pellet cells for 10 min at 6000 rpm (use SSA rotor).
3. Resuspend cells gently in 9.5 ml TE buffer. Add 0.5 ml of 10% SDS and 50  $\mu$ l of 20 mg/ml proteinase K. Mix thoroughly and incubate 1 hr at 37°C.
4. Add 1.0 ml of 5M NaCl and mix thoroughly.
5. Add 1.5 ml CTAB/NaCl solution. Mix thoroughly and incubate 20 min at 65°C. (can stay for a moment).
6. Add an equal volume (13 ml) of chloroform/isoamyl alcohol. Extract thoroughly. Spin 10 min at 7000 rpm (use S334 room temperature to separate phases).
7. Transfer aqueous supernatant to a fresh tube using a wide-bored pipet.
8. Add an equal volume phenol/chloroform/isoamyl alcohol. Extraction the equal volume. Repeat three times.
9. Add an equal volume of chloroform/isoamyl alcohol.
10. Add 0.6 volume isopropanol and mix. (-20°C overnight) (-20°C can help to precipitate.)

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2

Continue:

11. Transfer the precipitate to 1 ml of 70% ethanol in a fresh tube (just brown-red one), by hooking it on the end of a Pasteur pipet that has been bent (by heating it) and sealed.
12. Drying the pellet (just in the air). Remove ~~start~~ white and stringy DNA to the ~~4~~ 2nd TE buffer.
13. Do the electrophoresis of nucleic acid (there are a lot of DNA and RNA).
14. Use Rnase to digest the genomic DNA overnight.

## Genomic DNA (continue)

1. Use *EcoR* I to do restriction. put
- |               |            |
|---------------|------------|
| genomic DNA   | 27 $\mu$ l |
| buffer (H)    | 9 $\mu$ l  |
| <i>EcoR</i> I | 4 $\mu$ l  |

overnight.

Digest genomic DNA.

2. do the electrophoresis at 60 V

put 109  $\mu$ l digested DNA (90  $\mu$ l digested DNA  
19  $\mu$ l dye)

the result is not good.

Extracting Plasmid DNA of *E. coli*. (Qiagen Midi and Maxi Pmt.)

1. Inoculate a 5-ml LB medium with *E. coli* 37°C shake to afternoon
2. Inoculate 200ml medium. Grow at 37°C ~~for~~ overnight.

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## Extracting plasmid DNA. (continue)

3. Harvest the bacterial cells by centrifugation at 6000g for 15min at 4°C
4. Resuspend the bacterial pellet in 4ml of buffer P1.
5. Add 4ml of buffer P2, mix gently but thoroughly by inverting 4-6 times, and incubate at room temperature for 5 min.
6. Add 4ml of chilled buffer P3, mix immediately but gently by inverting 4-6 times, and incubate on ice for 15 min.
7. Centrifuge at 20000 g for 30 min at 4°C. Remove supernatant containing plasmid DNA promptly.
8. Re-centrifuge the supernatant at 20,000 xg for 15 min at 4°C. Remove supernatant containing plasmid DNA promptly. ~~Alternatively,~~  
~~the sample can be fill~~
9. Equilibrate a QIAGEN-tip 100 by applying 4ml Buffer QBT, and allow the column to empty by gravity flow.
10. Apply the supernatant from step 8 to the QIAGEN-tip and allow it to enter the resin by gravity flow.
11. Wash the QIAGEN-tip with 10ml (two times) buffer QC.
12. Elute DNA with 5 ml buffer QF.
13. precipitate DNA by adding 3.5 ml room-temperature isopropanol to the eluted DNA. Mix and Centrifuge immediately at <sup>13000 rpm</sup> ~~15000 g~~ for 30 min at 4°C. Carefully decant ~~the~~ the supernatant.
14. Wash DNA pellet with 2ml of room-temperature 70% ethanol, and centrifuge at <sup>13000 rpm</sup> ~~15000 g~~ for 10 min. Carefully decant the supernatant without

disturbing the ~~pette~~ pellet.

15. Air-dry the pellet for 5-10 min. and redissolve the DNA in a suitable volume of buffer  $\rightarrow$  (TE buffer  $\rightarrow 1/2$  ml).
16. Do electrophoresis to check it.  $\rightarrow$  to digest.  $\rightarrow$  next page

8/3/99

### High-efficiency transformation by electroporation

1. Inoculate a single colony of E. coli cells into 5 ml LB medium. Grow 5 hr. to overnight at 37°C with moderate shaking.
2. Inoculate 2 ml of the culture into 500 ml LB medium in a sterile flask. Grow at 37°C <sup>shaker</sup> shaking at in incubator. ~~to an OD<sub>600</sub> of 0.5-0.7~~ <sub>as soon as possible.</sub>
3. Chill cells in an ice-water bath 10 to 15 min and transfer to a prechilled 50 ml centrifuge bottle.
4. Centrifuge cells ~~20 min~~ <sup>10 min</sup> at ~~2000 rpm~~ <sup>8000</sup> 2°C.
5. Pour off supernatant and ~~resuspend~~ resuspend the pellet in ice-cold water. Add 25 ml ice-cold water and mix well. <sup>at 10 min 8000 rpm. repeat 2 times.</sup>
6. Centrifuge cells as in step 4. <sup>Throw away the supernatant to the bleach bottle including</sup>
7. Pour off supernatant immediately and resuspend the pellet by swirling in remaining liquid.
8. Add another ml ice-cold water, mix well, and centrifuge again in step 4.
9. Pour off supernatant immediately and resuspend the pellet by swirling in remaining liquid.
7. Put remaining liquid to the eppendorf tube. centrifuge it at 12000 rpm 5 min. pour off by using vacuum.
8. add 100 µl water to the tube. mix well. put it on ice.

(余九芳力管)

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4

6

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10. Set the electroporation apparatus to 2.5 kV, 25  $\mu$ F. Set the pulse controller to 200 or 400 ohms (Under the machine instruction)
11. Add 1  $\mu$ L plasmid DNA (PGIP704) <sup>and 20  $\mu$ L bacterial</sup> in <sup>to</sup> tubes containing fresh or thawed cells (on ice). Mix by tapping the tube or by swirling the cells with the pipettor.
12. Transfer the DNA and cells into a uvette that has been chilled 5 min on ice. Shake slightly to settle the cells to the bottom, and wipe the ice and water from the uvette with a Kimwipe <sup>between</sup>.
13. Place the uvette into the sample chamber.
14. Apply the pulse by pushing the button of flipping the switch.
15. Remove the uvette. <sup>LB broth immediately</sup> Immediately add <sup>it</sup> ~~300  $\mu$ L medium~~ and transfer to a stir culture tube <sup>including LB broth</sup> with a Pasteur pipet. Incubate ~~to~~ 60 min with moderate shaking <sup>37°C</sup>.
16. Plate aliquots of the transformation culture on LB plates containing antibiotics.

plasmid digest

PGP ~~the~~ 1  $\mu$ L  
 10x buffer 1  $\mu$ L  
 DnaeI 7.5  $\mu$ L  
 EcoR I 0.5  $\mu$ L

run 5  $\mu$ L on gel37°C 2 hr.  $\rightarrow$  65°C for 15 mindilution  $10^{-6}$  100  $\mu$ L to LB DAP $10^{-5}$   $10^{-2}$  100  $\mu$ L to LB DAP DAP

result

 $10^{-5}$   $10^{-2}$  zero 2-3  $\times 10^6$  $10^{-6}$  100 colonies

To electrophoresis.

1. 15 ~~ul~~ <sup>ml</sup> marker (1Kb) first line
2. 65 ~~ul~~ <sup>ml</sup> digested DNA + 13 ~~ul~~ <sup>ml</sup> 6x dye second line
3. cut out 7-10 kb gene.
4. elute into Bio-rad - promega.

Fragment Isolation Protocol from NuSieve Gel.

1. Place the electrophoresed gel on the UV transilluminator and view under long wave light. Locate the fragment of 7-10 kb and cut out the band using a glass cover slip and place the gel fragment into a 1.5ml microcentrifuge tube.

2. Heat to 65°C until gel slice melts
3. Add 3 volumes of BioRad DNA isolation binding buffer and for every 2 <sup>120 ~~ul~~ <sup>ul</sup></sup>  $\mu$ g of DNA add 5 <sup>5 ~~ul~~ <sup>ul</sup></sup>  $\mu$ l of matrix <sup>30 ~~ul~~ <sup>ul</sup></sup>  $\mu$ l matrix
4. Attach a clean 3cc syringe to a promega spin column and add the solution to it.
5. Push this slowly through the column (1 drop/3 sec)
6. Detach column from syringe and remove plunger.
7. Wash with 2mls of 95% isopropanol.
8. Spin at 12,000 rpm for 20 sec.
9. Dry Column in Hybridization oven for a few minutes <sup>65°C in</sup> at 65°C
10. Add 30-50 ~~ul~~ <sup>ul</sup> of water or TE. wait 1 min. and spin again at 12,000 rpm for 20 sec.
11. Run 10 ~~ul~~ <sup>ul</sup> of the eluant on a gel to check recovery.

5 MB

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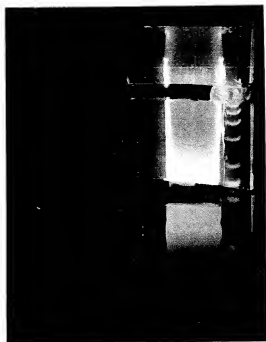
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8

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cut out 7-10 kb

6

5

4

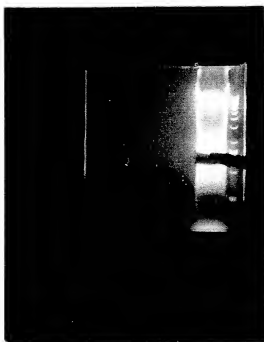
3

2

1

bright 1 kb

dig. tel. 0.5 craters (1 kb)



6

5

4

3

2

1

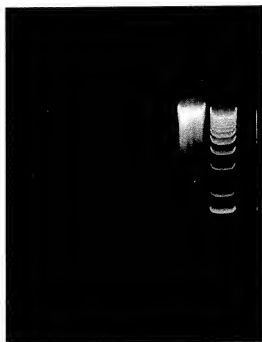
Do electrophoresis to check the result.

DNA. 5  $\mu$ l. + 1  $\mu$ l dye.

15  $\mu$ l Marker

$\rightarrow$  60v. run

Results:



Conjugation:

1. Donor Strain (*E. coli* MGN 617 PGP704 or MGN 617 PLOF Km) are grown overnight with shaking at 37°C in 2 ml LB broth containing 200  $\mu$ g/ml ampicillin and 50  $\mu$ g/ml kanamycin. (DAP)
2. Grown *P. multocida* 11039 overnight with shaking at 37°C.
3. 50  $\mu$ l MGN 617 or 50  $\mu$ l ~~11039~~ + 100  $\mu$ l 11039 to 5 ml of 10 mM  $MgSO_4$ .
4. Vortex for a few seconds.
5. Transfer to a 5 ml disposable syringe and filter through a 25 mm Nitric filter.
6. Drain the filter and carefully remove it from the filter case with sterile forceps (curved tip work best).

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10 100ul → 100ul

100ul 1PDR

Notebook Number:

200ul

Date:

08/05/99

7. Place the filter on a LB plate cell side up.  
(avoid bubbles between the filter and the agar)

8. Incubate at 37°C for 8-18 hrs.

PLOT

2.36V. Km

08/06/99

Results 9. Resuspend the filter using 3ml 10m MgSO<sub>4</sub>. Vortex.

PGP 704 BHI Ap 100ul 10<sup>0</sup>  
BHI ApK 100ul 10<sup>0</sup> } negative  
BHI 100ul 10<sup>-5</sup>

PLOT Km BHI ApKm 100ul 10<sup>0</sup> → negative  
BHI Km 100ul 10<sup>0</sup>  
BHI 100ul 10<sup>-5</sup>  
LB Ap Km DAP 100ul 10<sup>-5</sup>  
200ul

Transformation by electroporation

PGP 704.

PLOT Km 10<sup>0</sup> 10<sup>-1</sup> 10<sup>-2</sup> dilution

Results

08/09/99

Digestion.

93-146 genomic DNA (CL plasmid prep).

Leat incubate at 70°C 30min

Restriction Digestion EcoRI

DNA 51ul

10x buffer 6

EcoRI 3

37°C 4h → put in 65°C water bath

08/10/99

## 1b electrophoresis

9+3-146 genomic DNA (after digest) 60  $\mu$ ldye (6x) 10  $\mu$ l↓  
run gel (marker 1+6 12  $\mu$ l)

## Fragment Isolation Protocol from Gel (See Page 7)

1. Cut out 7-10 kb and place the gel fragment into microcentrifuge tube
2. Heat to 65°C until gel slice melts
3. Add 3 volumes (630  $\mu$ l) of BioRad DNA isolation binding buffer ~~and then~~ (1  $\mu$ g  $\approx$  1  $\mu$ l volume) and usually add 30  $\mu$ l matrix.
4. Attach a clean 3 cc syringe to a promega spin column and add the solution to it.
5. Push this slowly through the column.
6. Detach column from syringe and remove plunger.
7. Wash with 2 mls of 95% isopropanol.
8. Spin at 12,000 rpm for 20 sec.
9. Dry Column in Hybridization oven for a few minutes.
10. Add 30  $\mu$ l of TE wait 1 min and spin again at 12,000 rpm for 20 sec.
11. Run Gel.

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29  
9/30/99

Digest:

D1039 60ul

Buffer 7ul

EcoRI 3ul

37°C 2h / 65°C 15min

9/30/99

1. Run a gel. 70ul

2. Cut out 7-10kb.

3. Use another kit to extract DNA.

QIA quick Gel Extraction kit protocol (another kit)

1. Excise the DNA fragment from the agarose gel with a clean, sharp scalpel.
2. Weigh the gel slice in a tube. Add 3 volumes of Buffer QG to 1 volume of gel (100mg ~ 100ul)
3. Incubate at 50°C for 10min.
4. After the gel slice has dissolved completely, check that the color of mixture is yellow.
5. Add 1 gel volume of isopropanol to the sample and mix.
6. To bind DNA, pipet the sample onto the QIA quick column and apply vacuum. After the sample has passed through the column, switch off vacuum source.

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10/23/99

## Run gel (Southern Blotting)

1. Digest EcoRI SalI BamHI XbaI 11039EcoRI → in back
2. ~~Rinse & Rinse~~
3. photograph with a ruler laid alongside the gel so that band positions can later be identified on the membrane.
3. Rinse the gel in distilled water and place in a clean glass dish containing ~10 gel volumes of 0.25 M HCl. Shake slowly on a platform shaker for <sup>10 min</sup> ~~30 min~~ at room temperature.
4. Pour off the HCl and rinse the gel with distilled water. Add ~10 vol denaturation solution and shake as before for 30 min to 1 hr.
5. Pour off the denaturation solution and Rinse the gel with distilled water. Add ~10 vol neutralization solution. shake as before for 30 min to 1 hr.

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11/20/99

Conjugation:

See P10;

put the conjugation in

① MGN 617  $10^{-1}$  BHI Km②  $10^{-1}$  BHI Ap-③ MGN 617 PGP 704  $10^{-1}$  BHI ApKm- $10^{-1}$  BHI Km $10^{-1}$  BHI Ap $10^{-1}$  BHI ApKm

→ can grow



③ MGN 617 PLOF Km

 $10^{-1}$  BHI Km x2 (100 uL, 20-50 colonies) $10^{-1}$  BHI [Ap] $10^{-1}$  BHI ApKm $10^{-1}$  and  $10^{-5}$  BHI 11039 $10^{-5}$  LB DAP ApKm

→ MGN 617

↓  
200 uL DAP

11/23/99

Cut out 2-6 Kb.

7-10 Kb. and more than

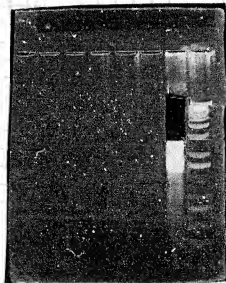
10 Kb

Use Qiagen gel

extraction kit to extract

DNA.

result:



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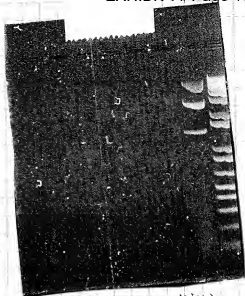
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the digest is not

in PTEM-32 has not been  
digested completely.

So there are three bands.  
(supercoiled DNA and one  
strand DNA)

So repeat the digest



PTEM-32  
plasmid DNA (1.4x)

11039 having 1 week lag can not grow in BHI broth.  
it can grow in BHI plate

Inoculate 11039 and 11039-617. PGM 704. PLOT km again

Do digest again 2.6 ul DNA (PTEM-32)  
1 ul EcoRI + 3 ul buffer  
over 2h. + 15' 65°C

1113/2000

Sap 1 ul + 4 ul buffer → 37°C 2h. → 15' 65°C

+ 20 ul 10% EtOH + 4 ul 3M NaAc → freeze

+ 20 ul 10% EtOH 13500 rpm 15' → pour off supernatant →

100 ul 20% EtOH 13500 rpm 10' → pour off supernatant

→ dry → run a gel (1 ul)

Conjugation LB plate + 200 ul ASP + 100 ul IPTG.  
do conjugation

BHI → BHI + 100 ul AP → BHI AP

BHI + 100 ul km → BHI km

BHI + 100 ul AP + 100 ul km → BHI AP km

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2/4/00

Run a gel. Digest. 21.5 ml DNA  
2.5 ml buffer overnight (37°C)  
1 ml EcoRI

→ 65°C 15'

Run a gel. Cut out 2-8 kb 6-10 kb. <sup>purified</sup>  
(because the extraction DNA always bigger than  
it is. so cut out <sup>the</sup> smaller band)  
<sub>little</sub>

2/7/00

Extract DNA. (use Qiaquick Spin <sup>kit</sup> ~~Handbook~~)

2/8/00

Run a gel. but no DNA

Clean DNA

improve: Clean DNA before

2/10/00

★ Clean DNA

1/10 volume 3M Na acetate

2 Vol 100% EtOH

-20°C ≥ 1h

Centrifuge ≥ 13,000 rpm 4°C 15'

Remove supernatant

Add 100% 100% EtOH

Centrifuge ≥ 13,000 rpm 4°C  
10 minRemove supernatant <sup>air</sup> dry ~ 5 minResuspend 30 µl DNase free H<sub>2</sub>O

Run a gel → cut out

2/13/00

1. Run a gel (genomic DNA 11039)

2. Do digest

just has a little DNA  
may because the DNA doesn't  
beet dissolve well.

3/11/00

Set up PCR reaction

	①	②	③	④
primer SC 1011	0.8		0.8	0.8
primer SC 1012	0.8		0.8	4
dNTP	4	4	0	8
3.3x buffer	6	6	6	6
Mg	2.4	2.4	2.4	2.4
H <sub>2</sub> O	<u>6</u>	<u>7.6</u>	<u>10</u>	<u>6</u>
	20ul	20ul	20ul	20ul
higher layer				
control DNA	0	0	0	0.5
3.3x buffer	9	9	9	9
enzyme	1	1	1	1
H <sub>2</sub> O	<u>20</u>	<u>20</u>	<u>20</u>	<u>19.5</u>
	30ul			

put 11039 in 50° for 2 hrs because a lot of  
DNA do not dissolve in TE buffer more than

Run a gel (11039 genomic DNA)

① line 1 ul DNA  
② line 3 ul DNA

Do digest (11039) 7.5 ul total

62.5 ul DNA

7.5 ul buffer

5 ul Enzyme

37° overnight

65° 15'



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in 3 sq. rubber maid dishes put 2 pcs chromatograph filter paper (this is too soak up buffers)

- 1) Denaturation solution - takes very little just enough to soak up filter. 5 min.
- 2) Neutralization buffer 5 min.
- 3) 2x SSC 5 min. place it on the paper and dry for 30 cross link on program c3 → put them in plastic.

4113100.

1. preheat the hybridization buffer to 42°C
2. In a suitable container prewet the blot in 5xSSC. Loosely roll the blot and place inside the tube. Add small amount of 5xSSC to the tube and "unroll" the blot ensuring no air bubbles are trapped between the membrane and the tube. Don't allow the blot to overlap itself
3. Pour off the 5xSSC and add the appropriate volume of ~~the~~ hybridization (20ul). buffer.
4. Prehybridize in oven for at least 30 min at 42°C
5. prepare the labeled nucleic acid probe as instruction P13.

- ① Dilute the DNA to be labelled to a concentration of 10 µg/ml using the water supplied.
- ② Denature 100 µg of the DNA sample (100 µl) by heating for 5 min in a boiling water bath. <sup>by 2 ul + 3 ul dam + 7 ul H<sub>2</sub>O</sup> (3 ul dam + 7 ul H<sub>2</sub>O)
- ③ Immediately cool the DNA on ice for 1 min. spin briefly in a microcentrifuge to collect the contents at the bottom of the tube.
- ④ Add an equivalent volume of DNA labelling reagent (100 µl) to the cooled DNA. mix gently but thoroughly

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5/14/00

prepare sequencing for genomic DNA

Set up sequence program. (Liang sequence)

Roe-Lab	Big dyc	2-3mer primer	purified gen DNA	H <sub>2</sub> O
1x reaction	8ul	(10mM)	1039 base	5ul
		1ul (30mM)		

total 20ul

p.s. primer (add 3ul primer to 97ul H<sub>2</sub>O)

HAPDAMP2

HAPDAMP1

PCR

5/2/00 prepare sequencing

## 1. Column Hydration

- ① Remove the top of column, then add 0.8ml of reagent grade water. <sup>invert for a few times</sup> make sure no bubbles. leave the column for at least 2 hrs. at room temp.

## ② 2. Removal of Interstitial Fluid:

- ① remove the top cap first, then remove the end stopper from bottom
- ② allow excess column fluid to drain into a wash tube discard this fluid.
- ③ spin the column and wash tube in centrifuge at 750g for 2 mins discard it.

## 3. Sample processing:

- ① Add 20ul <sup>sequence PCR</sup> ~~PCR~~ production into ~~wash~~ column. make sure it's be into the column, but don't touch the column. (it's on another tube)
- ② spin the column at 750g for 1 min - the purified sample will collect in the collection tube
- ③ Dry the sample in a vacuum centrifuge

## 4. vortex resuspend sample in 25 ul of template reagent in

- ① vortex 30sec } X2 2 min denature in 95°C
- quick spin } Chill 5 min

put 10ul in sequencing tube  
freeze remaining 15ul at -20°C

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H. sequencing didn't work

Signed

11800 ~~fold~~  $\Rightarrow$  178000 colonies / ml

$$\frac{1.78 \times 10^5}{0.005 \text{ ug}} = 3.56 \times 10^7 \text{ / ug DNA}$$

Result: the problem is xgal. the old xgal is not useful for producing blue colonies

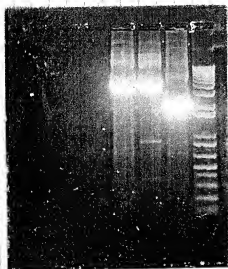
Do electroporation again, and inoculate the restreaked bacteria to the LB Broth.

6/13/00

Extract the plasmid DNA (restreaked Bact.)

put the electroporation to refrigerator

Run a gel (plasmid DNA # ① ② ③)



2ul + 3ul + 1ul dye  
DNA H<sub>2</sub>O

② and ③ is definitely inserted. do the digest.

20 ul. total

1 ul DNA

2 ul buffer

0.5 ELK I

16.5 ul H<sub>2</sub>O

overnight at 37°C load 3-4 ul on a gel.

6/14/00

Do electroporation. southern hybridization:

Run a gel (plasmid DNA ELK I digested. (20 ul))

load 3-4 ul per well.

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Date:

\* 2x SSC 5 min X2 at room temp.  
 5x SSC 0.5% SDS 42°C 30' 轻, oven  
 洗2遍 50 ml 20 ml 10%  
 2x SSC 用软纸 (Kimwipes) 将膜的正面擦。  
 3x SSC 洗干净 rinse and shake 5 min  
 5x 2

7/4/00

Do ligation

1 ul PGEM-32 EcoRI + Sap tt.  
 8 ul insertor (11039 3-5 kb extraction)  
 1 ul buffer  
 0.5 ul ligase (New England Lab)  
 16% overnight

7/10/00

65°C 15'

Do electroporation.

Note: cuvette should be chilled 5 min on ice  
 immediately add 1 ml LB medium and transfer  
 to a sterile culture tube.

12:100

Run a longer gel for 11039 E. coli.

1 lane: ladder (2ul) 5830 pm. 20V.

3 lanes: 11039 E. coli.

5 lanes: ladder (4ul)

It goes very slowly. I should run this gel at 30V.

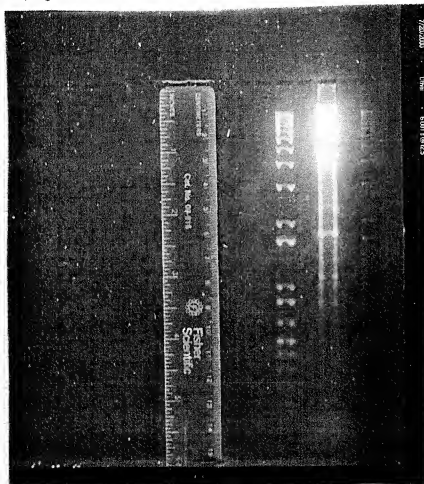
gel size 6x15

paper towel 8x15

filter paper 3 (55x14)

filter paper 5 (8x15)

membrane 7x14



the part from 6 kb to 8 kb has been separated. Do Southern hybridization.

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Make hybridization buffer.

100ml buffer

5 g blocking reagent

0.001M NaCl 2.925 g.

room temp. stir for 1 hr.

then 42°C at least 30'.

8/03/00

Continue Southern hybridization and Sequencing.

8/04/00

Sequencing didn't work again, but Southern hybridization did work.

Set up the PCR reaction to make more 1-10-2 and check the plasmid DNA.

primer dilution

45 ul H<sub>2</sub>O

2.5 ul (100 pmol/l)

25 ul --- p1

100 ul

→ 144.

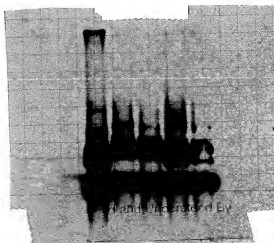
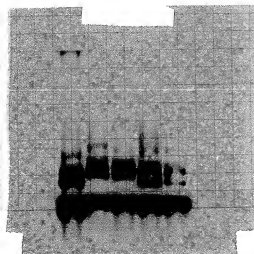
PCR reaction

① 1-10-2	0.5
dNTP	5
primers	1
buffer	5
tag	0.5
H <sub>2</sub> O	38.5

② plasmid	1 ul
dNTP	5
primer	1
buffer	5
tag	0.5
	37.5

③ 1-8-1	2
dNTP	5
primers	1
buffer	5
tag	0.5
H <sub>2</sub> O	36.5

④ Control	0
	5
	1
	5
	0.5
	38.5





Date: \_\_\_\_\_

8/25/00

Run a sequencing reaction

big dye	3 ul
plasmid	2 ul
primer	2 ul
H <sub>2</sub> O	8 ul

Should keep

~~some~~ a little

11039 Hind III

into

21

Left for Southern in a

✓ - region

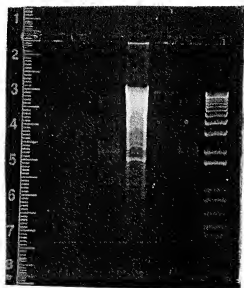
Run a gel for genomic DNA  
(5%) in order to <sup>comp</sup>sepe

8/28/00

Run a gel. 2ul for extracted DNA  
2ul for EcoRI 11039 digest (positive control)

- ① extracted DNA 4-6 kb
- ② 11039 EcoRI 6-9 kb
- ③ repeat ① → another tube
- ④ repeat ② → another tube

8/29/00



① ② ③ ④ ⑤

the size of both tubes is bigger.  
than needed. So extract DNA  
from 3-4 gel slice.

8/31/00

Run a gel

10/9/00

Digest - clone - a1 - a plasmid DNA to get probe

3ul  
digest

3 ul DNA

3ul buffer

1ul EcoRI ~~hmdIII~~ EcoRI23ul H<sub>2</sub>O

30ul → 2h. 37°C

15' 65°C

cut out from 1.4 kb.

pGEM-3Z hmdIII.

10/10/00

1. Continue plasmid DNA + sap treatment
2. do gel extraction.

10/11/00

Run a gel for gel extraction (for probe) and  
pGEM-3Z hmdIII + sap trt. (both are 5ul)  
gel extraction (probe) just need to use 3ul  
for detect.

I use PBS 1.5ul

Insert 7.5ul

buffer 1ul

Ligase 0.5ul

10/12/00

the electroporation didn't work. try  
electroporation.

1ul pGEM3Z + 10 ul LB

LB 10<sup>-2</sup>, 10<sup>-4</sup>.

10/13/00 it did work.

Read and Understood By

Signed

Signed

1/31/01

Do per again

	①	②	③
Pmltoxin A:3	0	1	1
dNTP	5	5	5
primer	1	1	0.5
buffer	5	5	5
New tag	0.5	0.5	0.5
H <sub>2</sub> O	38.5	37.5	38.0
	50 ul		

digest

DNA

buffer

EcoRI

H<sub>2</sub>O

①

②

③

1

1

1

5

5

5

0.5

0.5

0.5

43

43

43

50 ul

95°C 2 min

95°C 30 sec

72°C 10 min

4°C hold

64°C 1 min

72°C 2 min

1 cycle

35 cycles

2/01/01

Do PCR again. Add  
more genomic DNA and  
increase annealing  
~~decrease~~ annealing  
temperature to 66°C.

there're  
a little DNA on  
the gel. It's  
not clear. So  
Add more genomic  
DNA to do PCR  
again

	①	②	③	④
Pml (A:3)	0	3	6	10
dNTP	5	5	5	5
primer	1	1	1	1
buffer	5	5	5	5
Tag	0.5	0.5	0.5	0.5
H <sub>2</sub> O	38.5	35.5	32.5	28.5

50 ul



Q 120

annealing temperature 66°C 1'

Read and Understood By

Signed

Signed

3/5/01

Run a gel for H digest.

Do a Klenow for clone A ~~to~~ clai and T-H111  
digest

Klenow: 20- $\mu$ l digest (the concentration of my DNA is  
high enough)

0.8  $\mu$ l dNTPs (200 $\mu$ l of each in working  
solution)

1  $\mu$ l 0.1 mg/ml BSA (I skip this step)

0.5  $\mu$ l Klenow

I didn't add buffer because the Multi Core (buffer)  
is compatible for Klenow

incubate @ 37°C for 30 min. HI @ 75°C for 10 min.

(I lost this Klenow product)

3/6/01

Do digest for B. C. E. F. G. H. and Rnd a gel  
for it.

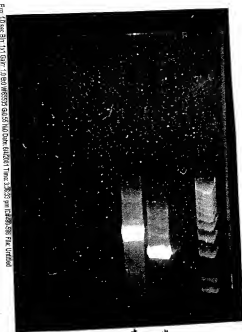
I select B. C. G. to run a seq. reaction again.

And to make sure which one is right.

Read and Understood By

Signed

Signed



PPS88 M13

6/5/01

Do the digest for PPS88

(Xba I + Kpn I)

5 ul DNA

12 ul H<sub>2</sub>O

(multi) buffer 2 ul

0.5 ul Xba I

0.5 ul Kpn I

2 h 37°C

inactivate 65°C 15'

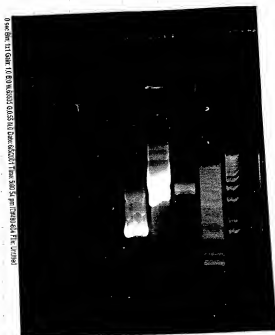
Run a gel for  
digestion, genomic DNA

PCLPM2

(from clone A)

and pbluescript plasmid

all the samples → 1 ul



PPS clone 88  
A plasmid

Read and Understood By

Signed

Signed

do sequencing react: for  
number 5

Run a gel for ① ② ③

big eye 8 μl  
plasmid 2 μl  
primer T3 2 μl

water 8 μl

Sample	Type	230nm	260nm	280nm	320nm	ratio	Concentration	Dilution	Factor	Purity
phagescript and dam	dsDNA	3.631	5.554	3.175	0	1.74929	0.2777 ug/ul	1	50	174.93

612901 number 5.

Sample	Type	230nm	260nm	280nm	320nm	ratio	Concentration	Dilution	Factor	Purity
pl 588	dsDNA	0.577	1.176	0.606	0	1.94059	0.0588 ug/ul	1	50	194.06

8 μl PLS 88

Xba I 0.5 μl

buffer (multicore) 1 μl

water 0.5 μl

8 μl PLS 88

Xba I 0.5 μl

Bam H I 0.5 μl

multicore 1 μl

7/3/01

PCR:

Template

pc1p m 3

0.5 μl

1 μl

control

dNTP

5 μl

5 μl

5 μl

primer

1 μl

1 μl

1 μl

buffer

5 μl

5 μl

5 μl

Tag

0.5 μl

0.5 μl

0.5 μl

H<sub>2</sub>O

38 μl

37.5 μl

38.5 μl

primer dilute

95 μl

H<sub>2</sub>O

25 μl

T7PI

25 μl

CL dam M20

588-8302-116	Jan 12 2001
GINOBYS	GINOBYS
TP1	TP1
5-ATACCTCCGCTTATACCGG	5-ATACCTCCGCTTATACCGG
TGAT	TGAT
588-8302-116	Jan 12 2001
GINOBYS	GINOBYS
TP1	TP1
5-ATACCTCCGCTTATACCGG	5-ATACCTCCGCTTATACCGG
TGAT	TGAT
588-8302-116	Jan 12 2001
GINOBYS	GINOBYS
TP1	TP1
5-ATACCTCCGCTTATACCGG	5-ATACCTCCGCTTATACCGG
TGAT	TGAT

588-8302-116	Jan 12 2001
GINOBYS	GINOBYS
TP1	TP1
5-ATACCTCCGCTTATACCGG	5-ATACCTCCGCTTATACCGG
TGAT	TGAT
588-8302-116	Jan 12 2001
GINOBYS	GINOBYS
TP1	TP1
5-ATACCTCCGCTTATACCGG	5-ATACCTCCGCTTATACCGG
TGAT	TGAT

GINOBYS

Read and Understood By

Signed

Signed

8/15/01 Cont.

PCR4 XLB electroporation...

Blotter plates with Hybond membrane  
followed protocol for colony hybridization

Cross-linked membrane -- stored at -20°C ON

8/16/01

Colony Hybridization PCR4 XLB

made probe using 3ul PCR-4 + 7ul H<sub>2</sub>O in ECL kit

preheat hybridization buffer to 42°C

wet screen mesh + membranes

Roll + put in hybridization tube

pre hybridize with 5X SSC for 1 hour

use 30mls hybridization buffer

probe Boil 5"

Cool on ice 5"

10ul labelling reagent

10ul glutaraldehyde

10° 37°C

add 1ml hybridization buffer from tube - add buffer +  
probe to tube 42°C ON

Friday

8/17/01

Colony Hybridization Cont.

preheat 1° wash buffer to 42°C

discard hybridization buffer. Add ~~5ml~~ 5ml

Add 50ml 5X SSC return to Oven 5min

discard 5X SSC add 1/3 vol of tube w/ 1° Wash Buf

Return to Oven for 20min

Wash again in 1° wash buff for 10min 2X

remove blots from tube place in glass container

Cover w/ 2X SSC

Shake @ RT for 5min X 2

Detection

8ml/membrane

incubate 1min

place on Saran

13mls #1

3mls #2

3membranes

Read and Understood By

Signed

Signed

Notebook Number: \_\_\_\_\_

Date: \_\_\_\_\_

9/19/01 Cont.

Culture reached 260 KU after 4 hrs incubation at 37°C

- (2) Put cultures into 534 tube
- (3) Incubate on ice 15"
- (4) Cf 4°C 5000 xg (6500rpm) 15"
- (5) Wash 2X w/ 1mM Hepes buffer (pH 7.0)
- (6) resuspend pellet in 10% glycerol to 1/12 their  
Original vol = 2 mls
- (7) Cf 5000xg 10"
- (8) resuspend in remaining glycerol solution after  
supernatant is decanted
- (9) Flash freeze in EtOH + dry ice bath
- (10) Freeze at -80°C

P.mut 11039 is highly encapsulated + forms a very soft pellet  
I lost the pellet in the 1st wash step, so more cells will  
have to be prepared

Electroporate PLS88 into P.mut 1069

1 µl PLS88 plasmid prep (in plasmid box sp# 99)

40 µl 1069 comp cells

2.5 KV  
200 Ω  
25 µF  
Tc = 5.12 msec

Incubate 37°C · 1 hr in LB

plate 10<sup>0</sup>, 10<sup>-1</sup>, 10<sup>+1</sup> 100 µl on LB strep 37°C

10<sup>+1</sup> = after plating, 10<sup>0</sup> + diluting 10<sup>-1</sup>, Cf culture + resuspend pellet in  
100 µl media - plate 100 µl

RESULTS

10 <sup>+1</sup>	134 colonies
10 <sup>0</sup>	2
10 <sup>-1</sup>	0

David and Hester 10/10/01



9/25/01 Tuesday

Pass colony from R6E12 #4 blot - do a plasmid prep  
Electroporate pLS88 into P.mult 11039

1  $\mu$ l pLS88 plasmid prep (in plasmid box spot 99)  
40  $\mu$ l 11039 comp cells (flash frozen in liquid N<sub>2</sub>)

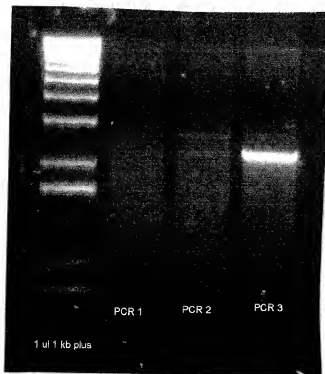
2.5KV  
200  $\Omega$   
25  $\mu$ F  
TC = 5.116 msec

incubate 37C 1 hr LB

Plate 10<sup>0</sup>, 10<sup>-1</sup>, 10<sup>+1</sup> 100  $\mu$ l on LB strep 37<sup>0</sup>C

RESULTS

10<sup>+1</sup> 80  
10<sup>0</sup> 13  
10<sup>-1</sup> 1



See pg. 35 for PCR rx setup

Annealing temp = 48<sup>0</sup>C

Extension time = 30 sec

PCR rx did not work  
again

0.7% agarose 3.5  $\mu$ l gel star  
5  $\mu$ l of PCR rx loaded

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Signed

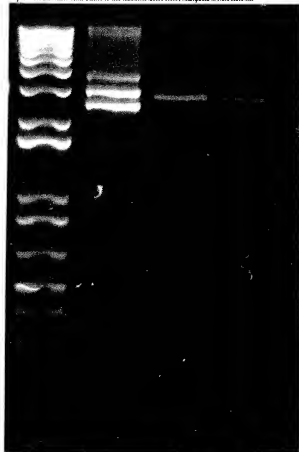
Notebook Number: \_\_\_\_\_

Date: \_\_\_\_\_

9/26/01 Wednesday - Gave to Memphis  
 ML started 5ml culture of ~~RoE12~~ <sup>RoE12</sup> Ed 9 +  
 9/27/01 Thursday

Did a Qia Spin mini prep of RoE12  
 Eluted in 30ul EB

Exp: 100 Sec 8.5 76359 0.416 Data: 01-40-1963 Time: 00:50 DNA: 00000000 File: 2591208 in 11039-25701.MF



1ul  
1kb  
Plus

RoE12  
1ul

PLS88  
in  
P.mult  
11039

RoE12 looks good  
 pass to a new plate  
 freeze next week

PLS88 in 11039 looks  
 good -

it work- we were  
 able to electroporate  
 into *P. multocida*  
 Comp cells



PCR pCLpm3 T7P1 + CLdamM20 092801

Sample #	1	2	3
pCLpm3 8/29/01 BYR	0.5	1	0
dNTPs	5	5	5
T7P1 + CLdamM20	1	1	1
Taq	0.5	0.5	0.1
Taq Buffer	5	5	5
Water	38	37.5	38
TOTAL Rx amt	50	50	
Annealing temp	67.5 C		

10/2/01



See pg 41 pcr  
Set up

pCLpm3  
1ul

1ul cut out of gel = 0.4177 gm  
put in Ref over right

10/3/01

Elute DNA from gel with Qiagen gel extraction Kit

cut gel fragment in 1/2 and put in 2 tubes

Added 600ul / tube of QG

Added 200ul isopropanol

eluted each column w/ 30ul EB into same tube = 60ul  
total

Called pCLpm3 PCR

Loaded 5ul of pCLpm3 PCR on 0.7% agarose gel 3.5ul gester  
100 Volts



Read and Understood By

Signed

Signed

Date: \_\_\_\_\_

10/3/01

Electroporated 40  $\mu$ l *P. multocida* 11039 Comp cells w/no DNA

Recover 1 hr in LB broth 37 C

plate 100  $\mu$ l 10<sup>0</sup>, 10<sup>-1</sup>, 10<sup>-2</sup> on LB

grow at 37 C O/N

Should have plated on LB strep -  
threw plates away2.5 K $\mu$ l25  $\mu$ l200  $\mu$ lT<sub>m</sub> = 5.16 min

Before Digesting pCLpm3 PCR 1.7 Kb frag we need  
more pcr product b/c XbaI has to be deactivated w/  
EDTA (heat does not work) + then the DNA is ppt  
out. DNA is lost in ppt so I need to amplify  
my PCR product

PCR pCLpm3 1.7 kb T7P1 + CLdamM20 10/3/01			
Sample #	1	2	3
pCLpm3 PCR 1.7 kb	0.5	1	0
dNTPs 1 mM each	5	5	5
T7P1 + CLdamM20	1	1	1
Taq (7/14/01)	0.5	0.5	0.5
Taq Buffer	5	5	5
Water	38	37.5	38.5
TOTAL Rx amt	50	50	50
Annealing temp	67.5 C		
extension time	30 sec		

run 0.5  $\mu$ l per rx on gel

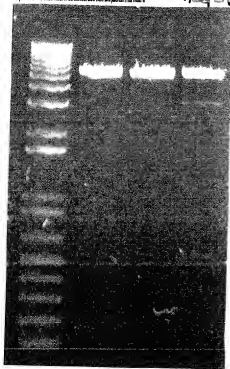
10ul PLS 88 plasmid prep 9/19/01  
1ul Xba I  
1ul Bam HI  
2ul multicore Buffer 10X  
6ul H<sub>2</sub>O

One

Digest 37C 4 hrs      Start 10:00am

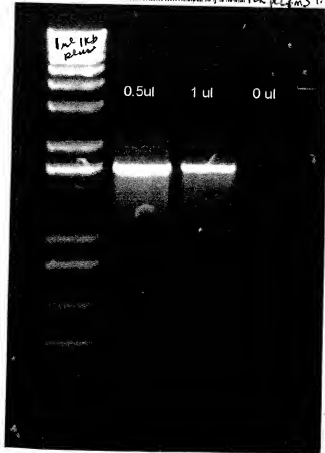
Run 0.5ul of pCp3.17Kb per Rx run 10/3/01 on  
0.7% agarose gel 3.5ul gel star

PLS88 Barnet & Baul Digest  
done 9/28/01



1ul 1ul 1ul 1ul  
1 kb Bam Kba Bam  
plus HE Kba I

Exp: 1930 Ser: B.0 W: 255 G: 1.65 Date: 00-00-1991 Time: 00:00 ID: 000000 File: C:\P40\17\104001.H PCR 1.7 Kb



Read and Understood By

Signed

Signed

Date: \_\_\_\_\_

10/4/01

Digest pCLPM3 1.7 Kb frag PCR Ex 1 (from 10/3/01)

0ul	DNA	
1ul	Xba I	
1ul	BamHI	37 C
2ul	MultiCore 10X Buffer	4 hrs
6ul	H <sub>2</sub> O	
<hr/> 20ul		

To pLS88 + pCLPM3 1.7 Kb BamHI XbaI digests

Add	0.5ul	0.5M EDTA	to deactivate Xba I
Add	2ul	3M NaAcetate	(1% vol)
	44ul	EtOH	<del>44</del> (2 vls)

freeze -20 over night to ppt DNA.

Start 2 1L BHKp cultures of ReE12 for LPS prep

Add 2.5mls of ON culture per Liter

Add 500ul 200mg/ml Ap = 100mg/ml final vol  
1/2 normal conc. used

grow ON 26°C Shaking

10/5/01

Cf. ReE12 cultures in 250ml bottles in C154 rotor

6000rpm for 15min at 4°C.

freeze pellets at -20°C.

Do LPS prep next wk

10/5/01

cf. ppet DNA pLS88 + pCLPM3 1.7kb  
resuspend in 10ul TE. put at 4C to go into  
suspension

10/8/01

run lal pLS88 + pCLPM3 1.7kb on gel  
✓ on gene spec first

Sample	Type	230nm	260nm	280nm	320nm	ratio	Concentration	Dilution	Factor	Purity
pCL PM 3 1.7 kb	dsDNA	0.655	1.536	0.818	0	1.87775	0.0768 ug/ul	1	50	104.32
	dsDNA	0.707	1.592	0.85	0	1.87294	0.0796 ug/ul	1	50	104.05
pLS88 Bam HI Xba I	dsDNA	1.566	3.982	2.369	0	1.68088	0.1991 ug/ul	1	50	93.38
	dsDNA	1.987	4.493	2.85	0	1.57649	0.22465 ug/ul	1	50	87.58

pCLpm3 76.8 ng/ul

pLS88 211.9 ng/ul

Ligation

pLS88 4.5 Kb

pCL PM 3 1.5 Kb

pLS88 1ul 200ng  
pCLPM3 7.5ul 570ng  
Ligase Buffer 1ul  
Ligase 0.5ul  
10ul

14°C overnight

10/9/01 Tuesday

Heat inactivate ligation 65°C 15"

Read and Understood By

Signed

Signed

## PCR 93146 EI OPS primers 10/17/01

Sample #	1	2	3	4
93146 genomic prep 7/28/99	2	2	2	2
dNTPs 2.5mM	10	10	10	10
EI OPS PCR U 2 + EI OPS PCR R 2	1.2	1.2	1.2	1.2
EXL polymerase	1	1	1	1
DMSO	0	0.5	1	1.5
Stabilizing Soln	1	1	1	1
10 X Buffer	5	5	5	5
Water	29.8	29.3	28.8	28.3
TOTAL Rx amt.	50	50	50	50
Use PCR EXL Pcr Program				
Annealing Temp	63			
PCR EXL primer				

Ran out of EXL polymerase  
after 1<sup>st</sup> rx.

1<sup>st</sup> rx  
ordered more EXL pcr

Rxs 2-4 are in green box  
- 20 chest freezer

gel picture pg 54

Make 2 5ml LBAP cultures of pFPV25 w/ DHI05s

- 1- plasmid prep
- 1- freeze

Make 5ml culture of pL588 isolates pos + Adj in LBstrep

positive colony  
from ~~best~~ adjacent to  
hybridization positive colony

Read and Understood By

Signed

Signed



Date: \_\_\_\_\_

10/18/01

Started cultures in wrong Antibiotics  
 started pFPVas w LB strep - pLs88 - LB Ap Opps -

Start new 5 ml cultures  
 pFPVas w LB Ap  
 pLs88 w LB strep

10/19/01

Froze 1 pFPVas culture (Ask B&K where it is)

Spin down other pFPVas culture + both pLs88 cultures  
 (pos + adj) - freeze pellets for plasmid prep

10/22/01

made Qiaspin plasmid prep of  
 PFPVas, pLs88 positive, pLs88 adjacent



Loaded 1ul of  
 pLs88 pos  
 pLs88 adj  
 pFPVas

Loaded 5ul of  
 PCR product

pass both pLs88 colonies  
 to a new LB strep plate

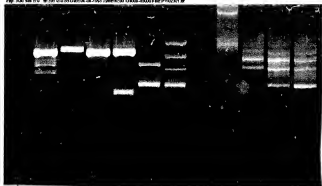
## Digest pLS88 pos + Adj plasmid prep (10/22/01)

Cuts out insert  
 pLS88 pos 4  $\mu$ l  
 Bam HI 1  $\mu$ l  
 Xba I 1  $\mu$ l  
 multicores 2  $\mu$ l  
 H<sub>2</sub>O 12  $\mu$ l  
 pLS88 Adj 4  $\mu$ l  
 Bam HI / Xba I cuts the insert out  
 20  $\mu$ l  
 load 10  $\mu$ l on gel

linearize  
 pLS88 pos 2  $\mu$ l  
 EcoRV 1  $\mu$ l  
 Buff D 1  $\mu$ l  
 H<sub>2</sub>O 6  $\mu$ l  
 10  $\mu$ l  
 pLS88 Adj 1  $\mu$ l  
 EcoRV 1  $\mu$ l  
 Buff D 1  $\mu$ l  
 H<sub>2</sub>O 7  $\mu$ l  
 10  $\mu$ l  
 load 10  $\mu$ l on gel

linearize DNA  
 pFPVas 5  $\mu$ l  
 EcoRV 1  $\mu$ l  
 Buff D 1  $\mu$ l  
 H<sub>2</sub>O 3  $\mu$ l  
 10  $\mu$ l  
 load 10  $\mu$ l on gel  
 pFPVas 5  $\mu$ l  
 pst 1  $\mu$ l  
 Buff H 1  $\mu$ l  
 H<sub>2</sub>O 3  $\mu$ l  
 10  $\mu$ l

Fig. 1000 Gel Doc 10/25/01 10/25/01 10/25/01 10/25/01 10/25/01 10/25/01 10/25/01



1 kb plus  
 did not show  
 up

- 1) pLS88 adj. Bam HI / Xba I  
10  $\mu$ l + 2  $\mu$ l 10x Dye
- 2) pLS88 pos EcoRV
- 3) pLS88 Adj EcoRV
- 4) pLS88 pos. Bam HI / Xba I
- 5) pFPVas pst I
- 6) pFPVas EcoRV
- 7) PCR 10/17/01 No DMSO 2  $\mu$ l DNA H<sub>2</sub>O 1  $\mu$ l + 3  $\mu$ l H<sub>2</sub>O
- 8) ↓ 0.5  $\mu$ l
- 9) ↓ 1  $\mu$ l
- 10) ↓ 1.5  $\mu$ l

Read and Understood By

Signed

Signed

Date: \_\_\_\_\_

10/24/01

PCR pMBE1 Ei OPS primers 10/24/01				
Sample #	1	2		
From 9/7/00 pMBE1 DNA box	0.5	0.5		
dNTPs 2.5mM	10	10		
Ei OPS PCR U 2 + Ei OPS PCR R 2	1.2	1.2		
EXL polymerase	1	1		
DMSO	0	0.5		
Stabilizing Soln	1	1		
10 X Buffer	5	5		
Water	31.3	30.8		
TOTAL Rx amt.	50	50		
Use PCR EXL Pcr Program				
Annealing Temp	63			
PCR EXL primer				

pL588 positive gene spec

Type	230nm	260nm	280nm	320nm	ratio	Concentration	Dilution	Factor	Purity
dsDNA	3.461	5.406	4.733	0	1.14219	0.2703 ug/ul	1	50	63.46
dsDNA	0.046	1.798	1.112	0	1.61691	0.0899 ug/ul	1	50	89.83
dsDNA	0.018	1.781	1.091	0	1.63245	0.08905 ug/ul	1	50	90.69
dsDNA	0.031	1.779	1.083	0	1.64266	0.08895 ug/ul	1	50	91.26
dsDNA	0.3	2.076	1.421	0	1.46094	0.1038 ug/ul	1	50	81.16
dsDNA	0.732	2.531	1.855	0	1.36442	0.12655 ug/ul	1	50	75.80
dsDNA	0.538	2.363	1.707	0	1.40187	0.11965 ug/ul	1	50	77.88
dsDNA	0.37	2.2	1.522	0	1.44547	0.11 ug/ul	1	50	80.30

pl588 plasmid prep made 10/24/01  $\text{Conc} = 124.8 \text{ ng/ul}$

Seq pl588 w/ CLPM1-PL 3.2pmol  $\rightarrow$  p, 29 bkr + w/T3 3.2pmol  $\rightarrow$  pg 27 bkr

PL588 4ul

TRR 8ul

primer 2ul

H<sub>2</sub>O 6ul

20ul

#1 CLPM1-PL

#2 T3

Sequence should contain DAM gene sequence

CLPM1-PL primer sequence data is trashy - but

I blasted a segment of it and it is the DAM gene sequence

Redigest pFPVas 5ul

Buff H

H<sub>2</sub>O

EcoRI

~~all~~

1ul

3ul

1ul

~~all~~

Pst I 1ul

10ul digest

pFP1as

EcoRI

Pst I

Buff H

H<sub>2</sub>O

5ul

1ul

1ul

2ul

10ul

20ul

Read and Understood By

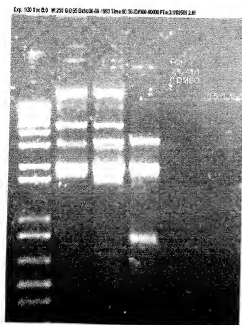
Signed

Signed

Notebook Number: \_\_\_\_\_

Date: \_\_\_\_\_

10/25/01



0.7% agarose 3.5 ul gel star

- 1) 1 ul 1 kb plus  
+ 4 ul H<sub>2</sub>O
- + 1 ul lax dyp
- 2) pFPV25- EcoRI 10 ul + 2 ul
- 3) pFA25 pstI 10 ul
- 4) pFPV25 EcoI/pstI 10 ul
- 5) PCR pmbEI 0.5 ul 20 DM
- 6 " " " 0.5 DMSC
- + 1 ul 6X dye
- 3.5 ul H<sub>2</sub>O

Freeze pl588 pos ~~XXXXXXXXXX~~  
Start 5ml LB strep

Electroporate pl588 pos into P. multi 11039 comp cells  
2.5 kV 40 ul P. multi (All D4 9/21/01)  
200  $\Omega$  1 ul pl588  
2.5  $\mu$ F  
TC = 5.08 ms Recover 1 hr LB strep

plate  $10^{11}$  on LB strep  
incubate 37°C ON

10/26/01

Exhibit A Page 46

Make Qiaspin plasmid prep of pFPV25 in DH10 $\beta$ s

Elute 30ul EB

Digest w/ EcoRI / Pst I (same as yesterday)

digest 5 hrs

run on gel w/ undigested plasmid

Freeze Pls ss now called pls ss DAM

it is in XL1B MRF'

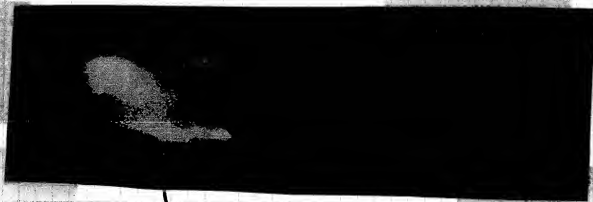
All D5-9 LB strep + 20% glycerol

Pick plate from electroporation in Ref Over weekend -  
Monday pick 4 big colonies + plate on LB strep

Started BHI Ap plate cultures of 93146 WT Lux +

93146 R<sup>-</sup> Lux

Imaged fish immersion  $\rightarrow$  1p injected w/ 93146 WT Lux  
on Night Owl.



$\rightarrow$  Luminescent area

Read and Understood By

Signed

Signed

pL588 positive gene spec

Type	230nm	260nm	280nm	320nm	ratio	Concentration	Dilution	Factor	Purity
dsDNA	3.461	5.406	4.733	0	1.14219	0.2703 ug/ul	1	50	63.46
dsDNA	0.046	1.798	1.112	0	1.61691	0.0899 ug/ul	1	50	89.83
dsDNA	0.018	1.781	1.091	0	1.63245	0.08905 ug/ul	1	50	90.69
dsDNA	0.031	1.779	1.083	0	1.64266	0.08895 ug/ul	1	50	91.26
dsDNA	0.3	2.076	1.421	0	1.46094	0.1038 ug/ul	1	50	81.16
dsDNA	0.732	2.531	1.855	0	1.36442	0.12655 ug/ul	1	50	75.80
dsDNA	0.538	2.363	1.707	0	1.40187	0.11965 ug/ul	1	50	77.88
dsDNA	0.37	2.2	1.522	0	1.44547	0.11 ug/ul	1	50	80.30

pL588 plasmid prep made 10/24/01  $\text{Conc} = 124.8 \text{ ng/ul}$ Seq pL588 w/ CLPM1-PL 3.2 pmol  $\rightarrow$  p, 29 bkr + w/T3 3.2 pmol  $\rightarrow$  pg 27 bkr

pL588 4ul

TRR 8ul

primer 2ul

H<sub>2</sub>O 6ul20ul

#1 CLPM1-PL

#2 T3

Sequence should contain DAM gene sequence

CLPM1-PL primer sequence data is trashy - but

I blasted a segment of it and it is the DAM gene sequence

Redigest pFPVas 5ul

Buff H

H<sub>2</sub>O

EcoRI

~~all~~

1ul

3ul

1ul

~~all~~

Pst I 1ul

10ul digest

pFPVas

EcoRI

Pst I

Buff H

H<sub>2</sub>O

5ul

1ul

1ul

2ul

10ul

20ul

Read and Understood By

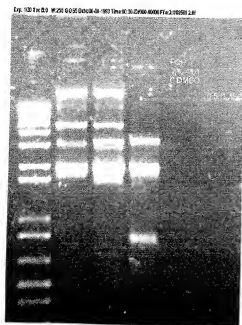
Signed

Signed

Notebook Number: \_\_\_\_\_

Date: \_\_\_\_\_

10/25/01



0.7% agarose 3.5ul gel star

- 1) 1ul 1 Kb plus  
+ 4ul H<sub>2</sub>O  
+ 1ul 6x dye
- 2) pFPV25- EcoRI 10ul + 2ul
- 3) pFA25 pstI 10ul
- 4) pFPV25 EcoI/pstI 10ul
- 5) PCR pmbE1 0.5ul 20 DM
- 6 " " " 0.5 DMSC  
+ 1ul 6x dye  
3.5ul H<sub>2</sub>O

Freeze pl588 pos ~~XXXXXXXXXX~~  
Start 5ml LB strep

Electroporate pl588 pos into P. multi 11039 comp cells  
2.5 KV 40ul P. multi (All D4 9/21/01)  
200  $\Omega$  1 ul pl588  
2.5  $\mu$ F  
TC = 5.08ms Recover 1 hr LB strep

plate  $10^{11}$  on LB strep  
incubate 37°C ON



10/26/01

Make Qiaspin plasmid prep of pFPV25 in DH10 $\beta$ s

Elute 30ul EB

Digest w/ EcoRI / Pst I (same as yesterday)

digest 5 hrs

run on gel w/ undigested plasmid

Freeze Pls ss now called pls ss DAM

it is in XL1B MRF'

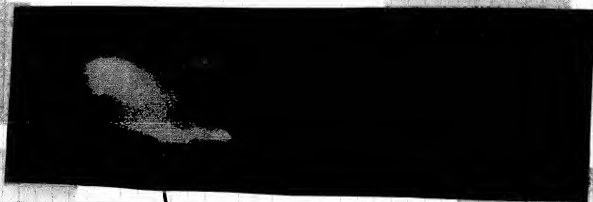
All D5-9 LB strep + 20% glycerol

Pick plate from electroporation in Ref Over weekend -  
Monday pick 4 big colonies + plate on LB strep

Started BHI Ap plate cultures of 93146 WT Lux +

93146 R<sup>-</sup> Lux

Imaged fish immersion + 1p injected w/ 93146 WT Lux  
on Night Owl.



→ Luminescent area

Read and Understood By

Signed

Signed

electroporation. grow on LB strep

11/5/01

pour 12%/4% SDS-PAGE gel  
gel stayed in buffer in 4°C ONStart 5ml LB strep cultures of P. mult 11039 pL588 From 10/31  
11/6/01

Samples to run on gel

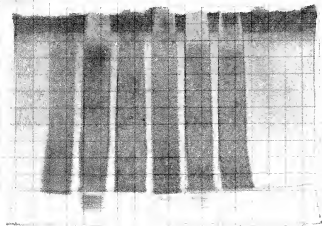
LPS from R6E12 dilute - 1/1 in sample Buffer

and take 10ul (1mg/ml) + 15ul H<sub>2</sub>O + 25ul Buffer = 10ugLPS Salmonella typh 1/1 of 2mg/ml in H<sub>2</sub>O 1/1 in Buffer

Lane 1	Sal typh	}	Not Denatured
2	R6E12 1/1		
3	R6E12 10ug		
4	Sal typh	}	Denatured
5	R6E12 1/1		
6	R6E12 10ug		

loaded 10ul / well Ran at 100 volts

Silver stain for LPS



Made Oligonucleotide plasmid prep of P.mult 11039 pL588  
+ colonies

11/7/01 Run 12% / 4% SDS PAGE gel

Same as 11/6/01

1) Sal. top  $\frac{1}{2}$  of 2mg/ml in H<sub>2</sub>O  $\frac{1}{2}$  in buffer - Load 10  $\mu$ l

2) R6E12  $\frac{1}{2}$  in buffer

3) R6E12 10  $\mu$ g

4) wt 93146 LPS 10  $\mu$ g

Electrophoresed at 100 volts

transferred to Nitrocellulose 100 volts 1 hr

Blocked membrane overnight 5% NFM in PBS

Our Ed 9 was contaminated - so was Bobbie's

DH103a pFPV25 frozen -80C All F1-5

tube w/1 used to start a 5ml LBap culture

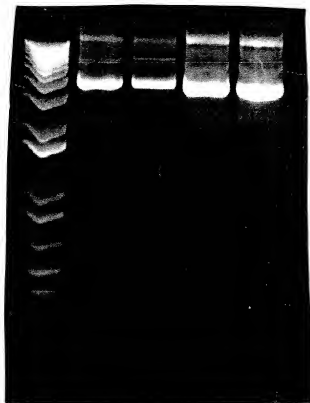
+ put on LBap plate

P.mult 11039 pL588 #1 Start 5ml LB strep culture to  
freeze  $\rightarrow$  see gel pic next page

Read and Understood By

Signed

Signed



p.mut 11039  
 Dan gene, pls 88  
 plasmid preps  
 from 11/6/01

1 ml  
 1K plus

11/8/01

Wester Blot R6E12

There is No ED 9 to use as 1<sup>st</sup> Ab so

- 1° used  $\alpha$  Ed 1c1 4383 polyclonal CF serum 1/1600  
 +  $\alpha$  Ed 1c1 93146 wt polyclonal CF serum 100ul added  
 to 1/10000 dilution

incubate RT 1 hr shaking  
 wash 3x PBS TW20

- 2° 9E1 1/4 1 hr RT shaking  
 wash 3x PBS TW

30 goat & mouse Ig Ap 1/1500 1 hr RT shaking  
 Develop BCIB/NBT 20 min

It did not work  
 will have to repeat  
 when Bobbie has more  
 Ed 9

1 2 3 4

Froze *pasturella multocida* 11039 pLSS8 (DAM gene)  
 LB + 20% glycerol  
 -80 C Box A11 G1-5

Digest pLSS8 p.mult 11039 #1

BamHI/XbaI

DNA 4  $\mu$ l  
 BamHI 1  $\mu$ l  
 XbaI 1  $\mu$ l  
 multicore 2  $\mu$ l  
 H<sub>2</sub>O 12  $\mu$ l  
 20  $\mu$ l

BAM HI

DNA 4  $\mu$ l  
 BamHI 1  $\mu$ l  
 multicore 1  $\mu$ l  
 H<sub>2</sub>O 4  $\mu$ l

Digest 37°C  
 for 2 hrs approx  
 run on gel tomorrow

Plasmid prep pFV95 in DH10 $\beta$ s  
 Eluted in 30  $\mu$ l set next day for gel pre

Read and Understood By

Signed

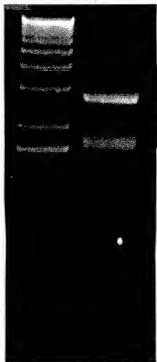
Signed

Notebook Number: \_\_\_\_\_

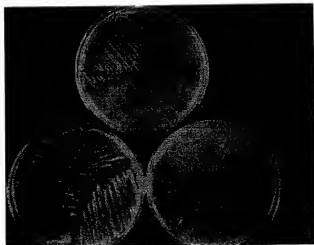
Date: \_\_\_\_\_

11/8/01

pFPR25



do not know what this band is

next run once and extract top  
band: electroporate into XL1B

Exp: 02 Sec: 01 Cam: 10.8.0 W: 630.0 H: 43.0 NO Date: 11/08/01 Time: 11:04:35 on Q640 file: 011111

Neg Controls  
14 + 15

22 = 93146 Lux

22 = 93146 Lux  
 22 = 93146 Lux  
 22 = 93146 Lux  
 22 = 93146 Lux

93146 Lux  
 recovered from  
 fish immersion  
 infected

Bacteria was  
 cultured  
 4 days  
 post  
 immersion in

Lux plasmid  
 is still  
 stable  
 in vivo

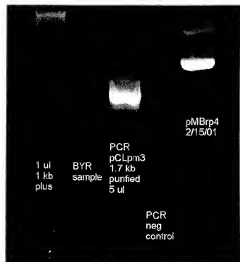
PCR pCLpm3 1.7 kb T7P1 + CLdamM20 purified  
01/16/02

Sample #	1	2	
pCLpm3 PCR 1.7 kb purified	0.5	0	
dNTPs 1mM each	5	5	
T7P1 + CLdamM20	1	1	
Taq (7/16/01)	0.5	0.5	
Taq Buffer	5	5	
Water	38	38.5	
TOTAL Rx amt.	50	50	
Annealing temp	67.5 C	program m1 under mark	
extension time	30 sec		

4/18/02

run Sml on gel of PCR rx

Sml DNA  
labeled by



measured pMBRp4 on gene spec - got  
a low reading of 11.8 ng/ul  
So rare lable on gel -  
Conc on gel looked like = 100 ng/ul

digest w/ BamHI HindIII

double digest  
pMBRp4 4 ul 4  
enzyme 0.5 ul 0.5 each  
10X Buff E 1 ul 1  
H2O 4.5 ul 4

digest from 2:00-5:00 37°C

Read and Understood By

Signed

Signed

2/13/02

Ligation: To End Conversion Rx mix add:

- (1) 1ul plasmid vector (pls & scale) digest  
1ul ligase
- (2) + control  
1ul pTZ Blue vector  
1ul ligase
- (3) - Control  
1ul pTZ Blue vector  
1ul ligase  
22°C 15"

~~plate~~

### Transformation:

Add 1ul ligation Rx. to 1 tube of Nova Blue Comp cells

Incubate on ice 5 min

heat shock 30 sec in 42°C water bath

Incubate on ice 2 min

Add 250ul RT SOC media

plate

(+) & (-) Controls 50ul @ 1/10 (5ul in 45ul H<sub>2</sub>O)  
on LB S-gal

exp. 50ul @ 1/10 on LB strep

2/14/02

Colonies from ligation were very small first thing  
this AM.

Read and Understood By

Signed

Signed



Date:

3/14/02 Thursday

PCR pCLpm3 1.7 kb T7Sal I P2 + CLDamECoRV 3/14/02			
Sample #	1		
pCLpm3 PCR 1.7 kb purified (1/23/02)	0.5		
dNTPs 1mM each	5		
T7Sal IP2 +CLDamECoRV	1		
Taq (7/16/01)	0.5		
Taq Buffer	5		
Water	38		
TOTAL Rx amt.	50		
Annealing temp	55		
extension time	30 sec		

used wrong template  
repeat using  
pCLpm3 plasmid prep  
8/29/01 red box 68



reconstituted primers

2.5ul each + 95ul H<sub>2</sub>O

made new 1mM dNTPs

40ml 10mM mix to 60ml H<sub>2</sub>O

plated plssss from frozen stock A3B4 put on LB strep at 37C. MLL tomorrow will take out of incubator + put at 4 C

Notebook Number: \_\_\_\_\_

Date: \_\_\_\_\_

4/17/02

Hybridize colonies from Missy's Pcr-p3 ligations  
 follow protocol pg 50

prime

pcr prod green pcr box A9 - pcr rx 3/21/02  
 pg 94 bkt - 535 ng/ul

Clean up pcr prod using microcon columns -  
 follow directs in kit

Removes primers + dNTPs

do head on gene spec -

Type	230nm	260nm	280nm	320nm	ratio	Concentration	Dilution	Factor
dsDNA	0.709	0.366	0.224	0	1.63393	0.0183 ug/ul	1	50
dsDNA	0.736	0.655	0.497	0	1.31791	0.03275 ug/ul	1	50
dsDNA	0.492	0.946	0.804	0	1.17537	0.04725 ug/ul	1	50
dsDNA	0.551	0.997	0.861	0	1.15796	0.04985 ug/ul	1	50

there are crazy readings - gene spec is  
 not great for measuring DNA of pcr prod.

MLL said use 3ul in probe

Remained of cleaned-up pcr prod is in Pink Box 118

Hybridize ON 42°C

4/17/02 Developed Hybridization

1" film almost had no image  
 10" very light image  
 1 fr OK

Plate 1

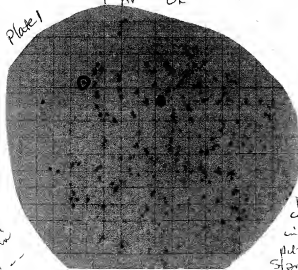
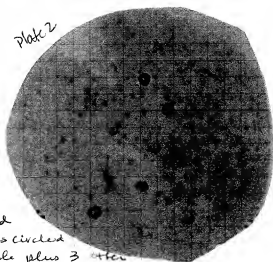


Plate 2



passed  
 colonies circled  
 in purple plus 3 others  
 put on LB strep  
 Start Sat 4/19/02 11:00 AM 11:00 AM 11:00 AM

plates in  
 1 -

5/13/02

cycle seq PGEM 3.2 test plasmid w/ M13 primer from Ab1  
 run on seq. to create matrix file

5/14/02 Tues

Clones from Electroporation grew well

grow 2ml LB strip cultures  
 pick colonies w/ sterile toothpick  
 use 1ml to transform to do plasmid preps for insert  
 grow 1 culture from Neg Control

plasmid prep	100ul	53
50ul	5	
Neg con (plasmid)	100ul	55
50ul	11	

pick 10 ligation colonies  
 1 neg con colony



5/15/02 Wed

all 2ml cultures grew

used 1ml to do Qimpreps spin plasmid preps

eluted in 30ul EB

Run 3ul DNA on gel  
 1ul GelDye  
 2ul H<sub>2</sub>O  
 6ul

It Finally  
 Worked !!

#5 & Neg  
 plasmid preps  
 in yellow box  
 A5 + A6

For Vaccine trial:

plated 93146 R<sub>6</sub> from 6/19/01 ATGS-80C on Blood  
 grow at 26°C 48hrs

Reconstituted Intervet Edict vaccine - filled bottle 2/3 full w/ sterile H<sub>2</sub>O

plate 1 loop full on Blood grow at 26°C 48hr

put 50ul into 5ml BHI to freeze back grow at 26°C shaking  
 Over night

1 750ul flask BHI for Intervet Vaccine  
 4 750ul flask BHI for R<sub>6</sub>

Read and Undersood By

Signed

Michael S.

Notebook Number: \_\_\_\_\_

Date: \_\_\_\_\_

For Vaccine trial cont:

Took 5mls of Heat Killed Ed 1ct (BYR's prep 4/12/02)

Tared tube empty

add 5mls

at 12000rpm 10"

Aspirate Sup

Re weigh tube = 0.0499 gm

Lyophilize overnight re weigh

resuspended pellet in 2mls H<sub>2</sub>OTransfer to a pre-weighed  
15 ml tube = 6.6018 gm

Freeze

Lyophilize

15 ml tube Dry wt = 6.6064

Dry wt of pellet = .0046

Digest pl<sup>+</sup> ss pel<sup>+</sup> m3 + clone (#5) + Neg plasmid preps

w/salt I + Cla I single digests

2ul DNA  
1ul Buffer  
0.5ul Enzyme  
6.5ul H<sub>2</sub>O  
10ul digest

ABI ~~matrix~~ matrix stds came in. stds are 4 colors: Blue, Red, Green, yellowmixed 1ul of each Standard with 12ul of deionized formamide  
(Borrowed from Hanson's Lab). 1 tube per std = 4 tubes 1 each colorHeat to 95C for 2"  
Chill on ice

Sequence using Seq Run (250ul) E run module

injected each sample 3 times - this is done when the injection  
list is filled out

Date: ~~11/10/02~~ 5/10/02

5/10/02 Thurs

Stds are still running on seq.

Weighed lyophilized pellet of heat ~~prepare~~ 2-750 ml cultures of heat killed Ed. 1ct - MondayRun digests on gel pL88 pol prn  $\checkmark$  electroporate plasmid prep clone #5

Load entire Digest

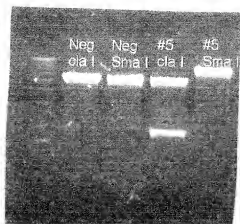
GK dye is really conc. So only use 100  $\mu$ l of 93146 #19 pass from MW plate10  $\mu$ l DNA digested1  $\mu$ l 6X Dye1  $\mu$ l 1+20

0.7% agy Make 2 more 750ml broth cultures

Run 100 Volts

12  $\mu$ l on gel

Lane 1 1  $\mu$ l 1Kb  
 2 Neg Cla I  
 3 Neg Sal I  
 4 #5 Cla I  
 5 #5 Sal I



Electroporate: pL88 into 11039 p.mult. electrocompetent cells

1  $\mu$ l plasmid prep #5 (yellow box) A8  
40  $\mu$ l 11039 comp cells box A11 D2

Tc = 5.12

5.5 KV

200  $\Omega$ 25  $\mu$ F

Recovered 1 hr in LB 37C

plate 100  $\mu$ l neat on 2 LB Strep plates

Made 2 more 750ml broth cultures to grow 93146 wt #19

to heat ~~kill~~ Kill for vaccine trial

Read and Understood By

Chickell S  
 Signed

379  $\times$  750ml = 27.75 gms BHI

5/16/02 Thurs.

Froze

pls Dam 2 5 Vials 1ml/each in LB glycerol + 30% glycerol  
Aguavac-Esc 5 Vials 1ml/each in BHI to 20% glycerol  
Vaccine

5/17/02 Fri:

Staff Appreciation Day

Nothing has grown yet on plates from Electroporation -

maybe need to be on BHI strep  
Maybe 11039 comp cells Dead

Leave at 37 C until Tomorrow

Passed 93146 WT #19 to new blood plate

Start 5ml culture Sun

Took R<sub>0</sub> + vaccine plates out of incubator  
leave at RT

Start 5ml BHI cultures Sat.

5/19/02 - Started 5ml BHI cultures

5/20/02 MM have Mark Sick BYA started 2 750ml cultures  
of ~~WT~~ WT #19

5/21/02 93146 WT Ed. 1st take 1ml from each 750ml culture  
Combine in 1 tube

Streak for purity - plate is pure

plate 10<sup>-5</sup> + 10<sup>-6</sup> for colony counts

10<sup>-5</sup> TNTC TNTC

10<sup>-6</sup> 286 287 Avg 286.5

Heat Kill at 60°C for 3 hours  
plate for Viability - NO growth oct 48 hrs

7/24/02

Blotted plates on Nitrocellulose membrane

Block in 0.01M PBS + 5% NFM 2 hrs RT

1° Ab used Straight Ed 9 1 hr RT

3X wash PBS 0.01M + Tween

2° Ab goat mouse Ig (H+L) AP 1/1500 40C Overnight

Set up Serum Killing Assay see pg 17 for plate setup  
Serum HI at 56°C for 30 min

Read plate every 5 min for 1 hr

plate 15 µl of  $10^{-4}$  +  $10^{-4}$  on BHI Ap

Colony Count Results (7/25/02)

E. coli  $10^{-4}$   
27, 30, 47 $10^{-7}$   
2, 2, 3assay worked  
per fecal  
 $2.31 \times 10^9$   
 $1.56 \times 10^9$ 

7/26/02 WT

 $10^{-6}$   
35, 21, 26 $10^{-7}$   
2, 5, 9 $1.82 \times 10^9$   
 $3.56 \times 10^9$ 

RC

 $10^{-6}$   
44, 30, 35 $10^{-7}$   
0, 3, 6 $2.42 \times 10^9$   
 $2 \times 10^9$ 

Injected mice

100 µl ip. of pinultocida WT 50, 100, 1000

2 mice per dose

" Dam 50, 100, 1000

" Dam 2 50, 100, 1000

Controls PBS 3 mice

Read and Understood By

  
 Signed

Signed

24 CO<sub>2</sub> indicates mouse was euthanized to  
reduce suffering

Notebook Number: \_\_\_\_\_

Date: \_\_\_\_\_

7/25/02 Mortality from mice Exp.

9:00am WT 1000 2 mice 1<sup>st</sup> DOA 2<sup>nd</sup> CO<sub>2</sub> lethargic, Shivering, Flaccid, piloerection  
WT 50 1 DOA  
Dam 1000 1 DOA  
Dam 100 1 DOA  
Dam 2 100 1 DOA

Notes:

2nd mouse in WT 50 tit is very lethargic, does not respond to being handled  
2nd mouse in Dam 1000 is lethargic, Shivering (due to fever?)

11:30am WT 100 CO<sub>2</sub> lethargic, Fever (shivering) does not respond to handling  
Dam 1000 1 DOA  
Dam 50 1 DOA

Notes:

2nd mouse in Dam 50 is sick - Shivering, non-responsive, piloerection

1:30 No more

4:45 WT 50 1 DOA  
Dam 50 1 DOA  
WT 100 1 DOA

All mice from Tit Groups WT 1000, WT 100, WT 50, Dam 1000, Dam 50  
are Dead

Dead mice were opened ventrally, the chest was spread open  
they were stored in Sample cups w/ 10% Buffered formalin

7/26/02

9:00am Dam 100 1 DOA

All mice from tit Group Dam 100 are Dead

7/27 = 7/28 Dan Scruggs ✓ mice

Mon 7/29/02

Start WT, RB + E. coli. Low cultures on Philip + Sine Philip both

Mice - Am all Tit groups are active + Alert  
pn " " " "



8/28/02

Colony counts- Serum Killing Assay E. coli Lux

$$\begin{array}{r}
 10^{-7} \quad \begin{array}{r} 5 \\ 5 \\ 5 \end{array} \\
 \hline
 1.6 \times 10^{10} \text{ CFU/ml}
 \end{array}$$

$$\begin{array}{r}
 10^{-6} \quad \begin{array}{r} 35 \\ 51 \\ 25 \end{array} \\
 \hline
 3.7 \times 10^{10} \text{ CFU/ml}
 \end{array}$$

8/29/02

Re lux + WT Lux Colony Count plates were  
Contaminated too badly to count

9/5/02 Inject mice

BYR prepared p.mutacida 11039 WT + Dan 2 for  
injections

1 extra mouse un.injected

5 mice / dose 9 treatments

20 - 4 doses WT 5, 10, 50, 100

80 - 4 doses Dan 2 10, 50, 100, 500

5 Control mice - PBS

Inject mice w/ 100ul each i.p.

use 1cc syringe with 27 1/2 G needle

mice were not set up properly - we (Dan Scruggs + I) had to

redistribute mice to 5/cage before injection - mice were Stressed!

injected mice 11:30am ✓ 1:30pm all OK ✓ 4:15pm all OK

Read and Understood By

*Michaela Sa*  
Signed

Signed

Notebook Number: \_\_\_\_\_

Date: \_\_\_\_\_

9/6/02

✓ mice 8:00 am

WT 5 - 3 mortalities

1 lethargic, shivering, unresponsive

1 almost dead completely unresponsive, unable to walk euthanize w/CO<sub>2</sub>

→ died before I could euthanize

WT 10 - 4 mort

1 sluggish but responsive

WT 50 3 mort

2 sluggish but responsive

WT 100 3 mort

2 sluggish but responsive have pilo erection

Dun2 500 All OK

100 all OK

50 all OK

10 all OK

PBS (-) Control All OK

✓ mice 10:45 am

WT 100 both mice are shivering, lethargic, very rapid breathing

✓ mice 2:45 pm

WT 100 breathing very rapidly unresponsive

WT 50 " "

WT 10 Very lethargic

WT 5 Sick but alert

Dun2 - all alert but not as alert as controls

✓ mice 6:20 pm

WT No change

Dun2 No change

PBS No change

10-4-03

Actinobacillus - plasmid electroporation

1. Removed 40  $\mu$ l of actinobacillus competent cells from microfuge tube and added to a clean tube.
2. Added 1  $\mu$ l of plasmid DNA to the same tube.
3. Placed suspension in electroporation cuvette.
4. Set electroporator settings.
5. Electroperated and got a time constant = 5.26.
6. Placed cells suspension in 1 ml BHI<sub>100</sub> + strep.
7. Incubated for 1 hour at 37°C.
8. Diluted culture to  $10^{-2}$ .
9. Plated 100  $\mu$ l of  $10^0$ ,  $10^{-1}$ ,  $10^{-2}$  on BHI<sub>100</sub> + strep plates.
10. Incubated plates at 37°C overnight.

Results

10-5-03

## Plate Count

$10^0$	33
$10^{-1}$	8
$10^{-2}$	0



1. Starter cultures of HB101 30-90 band electroporation for plasmid prep.
2. Starter cultures of 93446 wt, R<sub>6</sub>, and EC 11229 for MK micro.

### Microtiter boiling with conjugation mixtures

1. Killed bacteria in microtiter plate.
2. Resuspended in 100  $\mu$ l HBSS 0.9% with Ampicillin.
3. Removed 5  $\mu$ l from resuspended plate and added to 295  $\mu$ l HBSS 0.9%.
4. Removed 5  $\mu$ l from the dilution plate and added to 35  $\mu$ l HBSS 0.9% in white bottom assay plate.
5. Added an additional 40  $\mu$ l HBSS to central plate and 40  $\mu$ l normal serum to assay plate.
6. Incubated at RT for 1 hour.
7. Took readings on Dmax at Times 0, 30, and 60.
8. Analyzed data in Excel.
9. Highlighted results were going back (-80°C).

Results: See next page. (86)

1. 30-93 HB101 30-90 band plasmid prep.

1. Removed plasmid DNA with Oxygen kit.
2. Loaded 3  $\mu$ l on 7% agarose gel.
3. Ran at 100V for 1.5 hours.

Lane 1 - 1 kb ladder

Lane 2 - prep 1

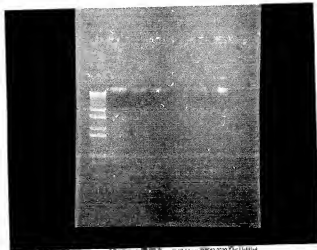
Lane 3 - prep 2

Lane 4 - prep 3

Lane 5 - prep 4

Lane 6 - prep 5

Lane 7 - prep 6



Exp: 1.8 sec Run: 1.5 hr Run: 100 V Run: 100 V Run: 100 V Run: 100 V Run: 100 V Run: 100 V Run: 100 V

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